SEQUENCE LISTING

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atgaggaaaa tgaaatgaat ggcacttatg actacagtca atatgaactg atctgtatca    180
aagaagatgt cagagaattt gcaaaagttt tcctccctgt attcctcaca atagttttcg    240
tcattggact tgcaggcaat tccatggtag tggcaattta tgcctattac aagaaacaga    300
gaaccaaaac agatgtgtac atcctgaatt tggctgtagc agatttactc cttctattca    360
ctctgccttt ttgggctgtt aatgcagttc atgggtgggt tttagggaaa ataatgtgca    420
aaataacttc agccttgtac acactaaact ttgtctctgg aatgcagttt ctggcttgta    480
tcagcataga cagatatgtg gcagtaacta aagtccccag ccaatcagga gtgggaaaac    540
catgctggat catctgtttc tgtgtctgga tggctgccat cttgctgagc ataccccagc    600
tggtttttta tacagtaaat gacaatgcta ggtgcattcc cattttcccc cgctacctag    660
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ccagctgcaa catgagcaaa cgcatggaca tcgccatcca agtcacagaa agcatcgcac    960
tctttcacag ctgcctcaac ccaatccttt atgttttat gggagcatct ttcaaaaact   1020
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ggttggggga aggggggagaa ataaaagcca agaagaggaa acaagataat aaatgtacaa   1320
aacatgaaaa ttaaaatgaa caatatagga aaataattgt aacaggcata agtgaataac   1380
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actctgctgt aacgaagaag agctttgtgg tgataatttt gtatcttggt tgcagtggtg    1440
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tttcaatttc ctggttttga cattatagta taattatgta agatggaacc attggggaaa    1560
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aaaacagatt attaaagagg ttcatgttaa aaggcattta taattatttt taattatcta    1740
agttttaata caagaacgat ttccctgcat aattttagta cttgaataag tatgcagcag    1800
aactccaact atcttttttc ctgtttttttt taaatttgta agtaatttta taaatccac    1860
ctcctccaaa aaagcaataa aaaaaaaaca aactataaaa aaaaaaaaaa aaaaaaaaaa    1920
aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaa                            1958
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<210> 2
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<212> PRT
<213> Homo sapiens

<400> 2
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Lys Glu Asp Val Arg Glu Phe Ala Lys Val Phe Leu Pro Val Phe Leu
            35                  40                  45
Thr Ile Val Phe Val Ile Gly Leu Ala Gly Asn Ser Met Val Val Ala
    50                  55                  60
Ile Tyr Ala Tyr Tyr Lys Lys Gln Arg Thr Lys Thr Asp Val Tyr Ile
65                  70                  75                  80
Leu Asn Leu Ala Val Ala Asp Leu Leu Leu Leu Phe Thr Leu Pro Phe
            85                  90                  95
Trp Ala Val Asn Ala Val His Gly Trp Val Leu Gly Lys Ile Met Cys
            100                 105                 110
Lys Ile Thr Ser Ala Leu Tyr Thr Leu Asn Phe Val Ser Gly Met Gln
    115                 120                 125
Phe Leu Ala Cys Ile Ser Ile Asp Arg Tyr Val Ala Val Thr Lys Val
    130                 135                 140
Pro Ser Gln Ser Gly Val Gly Lys Pro Cys Trp Ile Ile Cys Phe Cys
145                 150                 155                 160
Val Trp Met Ala Ala Ile Leu Leu Ser Ile Pro Gln Leu Val Phe Tyr
                165                 170                 175
Thr Val Asn Asp Asn Ala Arg Cys Ile Pro Ile Phe Pro Arg Tyr Leu
            180                 185                 190
Gly Thr Ser Met Lys Ala Leu Ile Gln Met Leu Glu Ile Cys Ile Gly
            195                 200                 205
Phe Val Val Pro Phe Leu Ile Met Gly Val Cys Tyr Phe Ile Thr Ala
    210                 215                 220
Arg Thr Leu Met Lys Met Pro Asn Ile Lys Ile Ser Arg Pro Leu Lys
225                 230                 235                 240
Val Leu Leu Thr Val Val Ile Val Phe Ile Val Thr Gln Leu Pro Tyr
                245                 250                 255
Asn Ile Val Lys Phe Cys Arg Ala Ile Asp Ile Ile Tyr Ser Leu Ile
            260                 265                 270
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Thr Ser Cys Asn Met Ser Lys Arg Met Asp Ile Ala Ile Gln Val Thr
        275             280             285
Glu Ser Ile Ala Leu Phe His Ser Cys Leu Asn Pro Ile Leu Tyr Val
        290             295             300
Phe Met Gly Ala Ser Phe Lys Asn Tyr Val Met Lys Val Ala Lys Lys
305             310             315             320
Tyr Gly Ser Trp Arg Arg Gln Arg Gln Ser Val Glu Glu Phe Pro Phe
            325             330             335
Asp Ser Glu Gly Pro Thr Glu Pro Thr Ser Thr Phe Ser Ile
            340         .       345             350


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<400> 3
actaccaaca ggttggtact tta                                        23


<210> 4
<211> 22
<212> DNA
<213> Homo sapiens

<400> 4
ctttgccatc tagagtggag cc                                         22


<210> 5
<211> 82
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<222> (1)...(82)
<223> n = A,T,C or G

<223> encodes synthetic peptide

<400> 5
ctttctattc tcactccgct gaannsnnsn nsnnsnnsnn snnsnnsnns nnsnnsnnsn    60
nsnnsccgcc tccacctcca cc                                          82


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<213> Artificial Sequence

<220>
<221> misc_feature
<222> (1)...(93)
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<223> n = inosine

<223> encodes synthetic peptide

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nnnnnnttca gcggagtgag aatagaaagg tac   93

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<220>
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<212> DNA
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<220>
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<220>
<223> mutagenic oligonucleotides

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<210> 10
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<212> DNA
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<220>
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<210> 12
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<220>
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aaagcgcagt ctctgaattt accg                                                    24

<210> 13
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<212> DNA
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<220>
<223> primers

<400> 13
tcgaaagcaa gctgataaac cg                                                      22

<210> 14
<211> 23
<212> DNA
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<220>
<223> primers

<400> 14
acagacagcc ctcatagtta gcg                                                     23
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